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OIPE

DATE: 01/30/2003 RAW SEQUENCE LISTING PATENT APPLICATION: US/10/053,498B TIME: 09:64:1 Input Set : A:\Seq Listing.txt Output Set: N:\CRF4\01302003\J053498B.raw 3 <110 > APPLICANT: Sloam-Kettering Institute for Canter Research Rothman, James E. Mayhow, Murk Hoe, Mer H. 7 Houghton, Alan Hartl, Ulrich Onerfoldi, Obathek Mersi, Yoichi 12 <120> TITLE OF INVENTION: CONJUCATE HEAT SHOCK PROTEIN-BINDING PEFTIDES 14 <130 > FILE REPERENCE: 11746/46002 18 KI40 CURRENT APPTICATION NUMBER: 10/053,498B 1' <141 CURRENT FILING DATE: 2002-01-17 19 <150 FRIOR APPLICATION NUMBER: 08/961,707 2) <151 PRIOR FILING DATE: 1997-10-31 22 K160 - NUMBER OF SEQ 10 NOS: 321 24 <170 GOFTWARE: WordFerfect 9.0 for Windows 76 HO10 SEQ ID NO: 1 H" KHILL LENGTH: 8 UB KO10 TYPE: PAT 29 021% OF GANISM: Artificial Sequence 31 -0.2 FEATURE: 53 4.25 OTHER INFORMATION: peptide in ml3 coliphage ::0' SEFUENCE: 1 · . . . His The Thr Val Tvr Gly Ala Gly ≥ 6 30 H. L. SEC ID NO: 2 40 -0.100 LENGTH: B 41 WILL TYPE: PRI 42 0.13 ORGANISM: Artificial Sequence 44 K. 200 FEATURE: 45 U.23 CTHER INFORMATION: peptide in ml3 coliphage 47 H400 REQUENCE: 2 The Glo Thr Ero Tyr Pro Thr Gly £(1) 50 -0110 SEQ ID NO: 3 53 0.110 LENGTH: 8 54 -0 110 TYPE: PRT 55 KW15: ORGANISM: Artificial Sequence 57 K2202 FEATURE: 88 <223> OTHER INFORMATION: peptide in m13 coliphage

60 <400> SEQUENCE: 3

62 63 Leu Thr Thr Fro Fhe Ser Ser Gly

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/053,498B

DATE: 11/av 2/03 TIME: 19:34:17

Input Set : A:\Seq Listing.txt
Cutout Set: N:\CRF4\01302003\J053498B.raw

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či klima sep ID NG: 4
66 <211> LENGTH: 8
67 KANAS TYPE: PRI
68 K213: CRGANISM: Artificial Sequence
PO KIRON FEATURE:
01 <223) OTHER INFORMATION: peptide in mli coliphage
73 ≈400: SEQUENCE: 4
(1) (ly Vai Fro Leu Thr Met Asp Gly
ī +
7: A210: SEQ ID NO: 5
78 4211) LENGTH: 6
80 K218: TYDE: FRT
81 <213: ORGANISM: Artificial Sequence
85 <2100 FEATURE:
$4 K213: OTHER INFORMATION: perside it #13 doliphage
BY KALON MEQUENCE: 5
   ...y. Let Pro Thr Val Let And Cly
50
84
91 (200) SEC ID NO: 6
3: <2011 LENGTH: 8
95 <2125 TYPE: PRT
9; <213. OFGANISM: Artificial Sequence
96 <210: FEATURE:
9" <2:3: OTHER INFORMATION: peptide in ml3 coliphage
9 + <400): SEQUENCE: 6
101 Cys Arg Phe His Gly Asn Arg Gly
102
104 00100 SEQ IE NO: 7
105 -0010: LENGTH: 8
106 -CHILLY TYPE: PRT
107 -001 -0 OFFANISM: Artificial Sequence
109 HOLDE FEATURE:
11: : :: :: OTHER INFORMATION: peptide in m13 coliphage
ILL JOHN SEPUENCE: 7
11.5
    Time Thr Arg Asp Phe Glu Ala Gly
115
117 H2100 SEQ ID NO: 8
11: *::11: LENGTE: 8
110 -011 - TYPE: PRT
1.13 4.21 a ORGANISM: Artificial Sequence
1.12 -1.1200 FEATURE:
123 4223 OTHER INFORMATION: peptide in m13 coliphage
1.33 <4000 SEQUENCE: 8
1.17 See Ser Ala Ala Gly Pro Arg Gly
1.13
130 JB1 JF SEQ 1D NO: 9
131 *211 * LENGTH: 8
132 KRIPW TYPE: PRT
133 <213> ORGANISM: Artificial Sequence
```

DATE: 1935 alls

TIME: U9:34:17

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Input Set : A:\Seq Listing.txt
                     Cutput Set: N:\CRF4\01302003\J053498B.raw
     135 KPZ > FEATURE:
     136 <223 - OTHER INFORMATION: peptide in ml3 ocliphage
     136 H400 H SEQUENCE: M
         Se: Let Ile Glm Tyr Ser Arg Gly
     143 KA10 - SEQ ID MO: 10
     144 <211 - LENGTH: 6
     148 <212 - TYPE: FRT
     146 K215 - ORGANISM: Artificial Sequence
     148 <220 PEACURE:
     149 <223 - OTHER INFORMATION: peptide in ml3 coliphage
     151 (12) FENTURE:
     15? <221 NAME/HEY: MISC FEATURE
     153 0222 LOCATION: 7)..(7)
     154 (123 OTHER INFORMATION: unknown
     155 (41) CEQUATAR: 18
W--> 158
         Asp Ala Leu Met Trp Pro Xaa Gly
     1 5 9
     161 (210 · SEQ ID NO: 11
     162 - 211 | LEMGTH: 8
     163 ARIZ TYPE: PRT
    164 (213 DEGAMISM: Artificial Sequence
    166 .22) FEATURE:
    16' 4883 - DIHER INFORMATION: peptide in m13 coliphage
    169 (22) · FEATURE:
    170 KOOL MAME, MEY: MISC FEATURE
    171 Chill LOCATION: (3)..(3)
    172 ROOTHER INFORMATION: unknown
    174 HADDLE SEQUENCE: 11
W--> 176
         Ser Ser Xaa Ser Leu Tyr Ile Gly
    1 "
    174 - 134 FED TE NO: 12
    130 -: 1 LENGTH: 8
     15° 4 1.6 TYPE: PRT
    182 - 133 OF GANISM: Artificial Sequence
    184 -0.000 FEATURE:
    135 - 17.30 OTHER INFORMATION: peptide in m13 coliphage
    180 H400H SEQUENCE: 12
    189 Phe Asr. Thr Ser Thr Arg Thr Gly
    1 10
    1 M -0 100 SHQ ID NO: 13
    193 <1110 IENGTE: 8
    134 KLIFF TYPE: PRT
    135 -0.180 CEGANISM: Artificial Sequence
    137 KILL FEATURE:
    198 30000 OTHER INFORMATION: peptide in m13 coliphage
    200 K400% SEQUENCE: 13
    202
         Thr Val Cin His Val Ala The Cly
    203
```

RAW SEQUENCE LISTING

FATENT APPLICATION: US/10/053,498B

1ATF: 1 % 2, 3

TIME: [3:34:1 FATENT APPLICATION: US/10/053,498B Input Set : A:\Seq Listing.txt Cutput Pet: N:\CRF4\01302003\J053498B.raw 278 921 > SEQ II NY: 14 206 <211> LENGTH: 8 ZUT <212: TYFE: PET 208 K213, ORGANISM: Artificial Sequence 110 k320 FEATURE: 311 k123. OTHER INFORMATION: peptide in m19 coliphage 113 (400) SEQUENCE: 14 215 As: Tyr Ser Phe Pro Pro Leu Gly 113 kg16, 8EQ IP NO: 18 319 <331 LEMOTH: 8 123 <232 TYPE: PAT 221 (213) ORGANISM: Artificial Sequence 123 THREE FEATURE: 221 CART OTHER INFORMATION: perhide in mlb coliphade CZOLIT CROMBET: This riy Her Mot Glu Her Let Gly 11.20 23. EBI KHOU FED ID NO: 16 230 CATO DENGTH: 8 133 Mil. TYPE: ERT 234 KEIN ORGANISM: Axtificial Sequence 336 KREET FEATURE: 237 KLES OTHER INFORMATION: peptide in m13 coliphage 239 Finds FEATURE: 240 KINIO NAME/KEY: MISC_FEATURE 241 HULDER LOCATION: (2)..(2) 24% RANGE OTHER INFORMATION: unknown 244 KODER FEATURE: 245 - 2222 NAME, KEY: MISC FEATURE 345 -0322 LOCATION: 36)..(6) HAV WILL OTHER INFORMATION: unknown ран сашин эторност: 16 W--> 251Phe Xaa Pro Met Ile Xaa Ser Gly 3.5.2 254 KHOW SEQ ID NO: 27 288 HUTTH LENGTH: 8 286 -271 - TYPE: PRT MSW 40.130 OF MANISM: Artificial Sequence HE9 HOLDE FEATURE: 1960 - CLIB OTHER INFORMATION: peptide in m13 coliphage HAR HIGORN SEQUENCE: 17 1004 Ala Pro Pro Arg Val Thr Met Gly . . . 5 5 0.67 ± 0.100 SEQ ID NO: 18 1168 -1111 LENGTH: 8 269 W. 12: TYPE: PRT 270 < 13: ORGANISM: Artificial Sequence 272 < 200 FEATURE:

RAW SEQUENCE LISTING

RAW SEQUENCE LISTING

FATERT AFFILICATION: US/10/053,498B

TIME: "A:54:1"

Input Set : A:\Seq Listing.txt
Set: N:\CRF4\01302003\J053498B.raw

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173 &183 > OTHER INFORMATION: peptide in mil coliphage 278 <400 > SEQUENCE: 18
     lie Ala Thr Lys Thr Fro Lys Gly
ZeC %ZICZ SEQ ID NO: 89
281 ×211> LENGTH: 8
282 K2122 TYPE: PRI
283 <2135 ORGANISM: Antificial Sequence
295 <220 · FEATURE:
286 K223 - OTHER IMPORMATION: poptide in mi3 belighade
Dark RANG - SEQUENCE: 12
23) Ty: Pro Pro Ten Phe Glm Tie Gly
231
293 <210 SEQ ID NO: 40
294 <211 LEMGTH: 5
기계 하다는 IMPH: PRI
196 <113 DESANISM: Artificial Sequence
299 :230 FEATURE:
23: 42:13 DIMER INFORMATION: peptide in mlb coliphage
301 -400 SEQUENCE: 20
3); Ty: His Thr Ala His Ash Met Gly
304
306 (210 SEP 10 NO: ..1
30' <211 - LEMGTH: 3
304 <212 - FYPE: PRT
3) + 213 - OEGANISM: Artificial Sequence
311 4000 FEATURE:
312 4003 OTHER INFORMATION: peptide in m13 coliphage
314 -4000 SEQUENCE: 21
316 Ser Tyr Ile Glr. Ala Thr His Gly
31.
519 d 160 PEQ 31 NO: 22
3.2 for 110 TabAGME: 3
BRI. C. . . INTER PRI
322 KLIB OFGANISM: Artificial Sequence
324 H. D. FHATURE:
325 K. 13: DTHER INFORMATION: peptide in m13 coliphage
327 RADGE SEQUENCE: 22
32+ Ser Fee Ala Thr Phe Leu Gly
330
337 -0410 SEQ ID NO: 23
334 KOLER TYPE: PRT
335 0.130 OFGANISM: Artificial Sequence
33' 40100 FEATURE:
53% CLESV OTHER INFOFMATION: peptide in m13 coliphage
340 <400> SEQUENCE: 23
312
   Thr Thr Pro Tro Aon The Ala Cly
343
```

RAW SEQUENCE LISTING ERROR SUMMARY FATENT AFFLI MITTON: US/10/053,498B

Input Set : A:\Seq Listing.txt

1 : run Net: N:\CRF4\01302003\J053498B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Con#:11; Maa Fis. '
Sen#:11; Maa Fis. '
Sen#:16; Maa Fos. 1,6
Sen#:47; N Fos. In
Sen#:46; N Fos. 6,10,1%
Sen#:56; N Fos. 6,10,1%
Sen#:130; Maa Fos. 6
Sen#:251; Maa Fos. 7
Sen#:251; Maa Fos. 7
Sen#:254; Maa Fos. 7